Chapter 2 Long-Term Vaccination and Treatment Strategies for COVID-19 Disease and Future Coronavirus Pandemics

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Abstract The appearance of new severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) variants with increased infectivity and immune escape capabilities has allowed continuation of the COVID-19 pandemic for the foreseeable future. This review describes the worldwide efforts aimed at developing new vaccination and treatment strategies to keep pace with these variants as they emerge. In the case of vaccines and monoclonal antibody-based therapeutics, we describe the development of variant-specifc, multivalent, and universal coronavirus directed approaches.

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Existing treatment approaches consist of repurposed medicines, such as antiviral compounds and anti-infammatory agents, although efforts are underway to develop new ways of preventing or minimizing the effects of infection with the use of small molecules that disrupt binding the SARS-CoV-2 virus to host cells. Finally, we discuss the preclinical and clinical testing of natural products from medicinal herbs and spices, which have demonstrated anti-infammatory and antiviral properties and therefore show potential as novel and safe COVID-19 treatment approaches.

Keywords Vaccination · COVID-19 · SARS-CoV-2 · Spike protein · Variant · Omicron

1 Introduction

As of June 23, 2022, 66.4% of the world population had received one or more doses of a World Health Organization (WHO)-approved COVID-19 vaccine, and over 12 billion doses have been administered in total [\[1](#page-16-0)]. However, the unequal distribution of vaccines has led to considerable moral outrage and could lead to epidemiological and economic disasters, as less than 20% of people in some low countries have received only one dose [\[2](#page-16-1)]. To compound the problem, the existing vaccines created to combat the original severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) strain which originated in Wuhan, China, may not work as effectively, if at all, against some of the newer SARS-CoV-2 versions, such as the Omicron subvariants [[3\]](#page-16-2). Despite the devastating effects the COVID-19 pandemic has had on our world, the imbalance in vaccination has still not been corrected, and there is still a signifcant proportion of the population in many countries and territories that show vaccine hesitancy [[4\]](#page-16-3). Thus, more studies are needed to understand and effectively correct this nonacceptance trend, which may threaten further efforts aimed at controlling the ongoing pandemic. Addressing the problem of how vaccines keep pace with new variants may be an even more diffcult prospect. In terms of keeping pace with the emerging variants, it is still not clear whether the best strategy is to develop vaccines against each variant as these emerge in a continuous game of catch-up, or if the construction of vaccines targeting multiple variants simultaneously is the best approach [[5\]](#page-16-4).

In the meantime, effective therapeutics may be needed for those individuals who are not fully protected by vaccination, or those who are immunocompromised or have a high risk of experiencing a severe COVID-19 disease outcome [[6\]](#page-16-5). Various monoclonal antibodies have been developed which target the SARS-CoV-2 spike protein, and some of these have demonstrated effcacy against the virus [[7\]](#page-16-6). However, as with the vaccines, many of these are only partially effective or completely inactive against some of the variants [[7\]](#page-16-6). The SARS-CoV-2 Omicron variant (lineage B.1.1.529) was detected in Botswana and South Africa in November 2021, and this spreads rapidly across South Africa and most of the world within 3 months

Fig. 2.1 Mutations of SARS-CoV-2 spike protein in the Omicron (B.1.1.529) BA.1 (top) and BA.2 (bottom) subvariants. The BA.1 subvariant contains 34 mutations and the BA.2 subvariant contains 31 mutations. Amino acid codes: A = alanine, D = aspartate, $E =$ glutamate, $F =$ phenylalanine, $G =$ glycine, $H =$ histadine, $I =$ isoleucine, $K =$ lysine, $L =$ leucine, $N =$ asparagine, $P =$ proline, $Q =$ glutamine, $R =$ arginine, $S =$ serine, $T =$ threonine, $V =$ valine, $Y =$ tyrosine

[\[8](#page-16-7)]. This rapid spread is likely to be due to the increased transmissibility and strong ability of this variant to escape immune detection by neutralizing antibodies [[9\]](#page-17-0). The property of increased transmission of this variant has been attributed to the enhanced capability of host infection via stronger interactions with the angiotensinconverting enzyme 2 (ACE2) receptor. The immune escape characteristic is a likely consequence of the higher number of mutations compared with other SARS-CoV-2 strains, rendering Omicron less recognizable to the existing vaccines and to convalescent sera from those had been infected by earlier strains (Fig. [2.1\)](#page-2-0) [[10,](#page-17-1) [11\]](#page-17-2).

This review describes the effects that the continuous variation in the SARS-CoV-2 genome has had on the efficacy of existing vaccines and treatments. This has created an urgent need to fne tune and advance new vaccine and drug development strategies to cope with this protein virus and to prepare for the next pandemic.

2 Current COVID-19 Vaccines

The current vaccines approved by the WHO are indicated in Table [2.1.](#page-3-0) These are based on different strategies which can be classifed as mRNA (Fig. [2.2a\)](#page-4-0), nonreplicating viral vector (Fig. [2.2b\)](#page-4-0), inactivated (Fig. [2.2c](#page-4-0)), and recombinant protein nanoparticle (Fig. [2.2d](#page-4-0)) vaccines. The frst of these to be approved by the WHO on Dec 21, 2020, was originally designated BNT162b1 and produced by Pfizer/ BioNTech [[12](#page-17-3)]. The International Non-proprietary Name (INN) is Tozinameran, and it is now sold under the tradename Comirnaty®. This was followed by Vaxveria (Oxford/AstraZeneca) [[13\]](#page-17-4), Covishield (Serum Institute of India) [\[14](#page-17-5)], Spikevax (Moderna) [\[15](#page-17-6)], Covilo (Sinopharm) [[16\]](#page-17-7), Ad26.COV2.S (Janssen) [[17\]](#page-17-8), and

		Countries approved		
Vaccine	Institution	(N ₀)	Approval date	Description
Comirnaty	Pfizer/ BioNTech	146	Dec 31, 2020	mRNA encoding spike protein
Vaxzevria	Oxford/ AstraZeneca	140	Feb 15, 2021	Non-replicating viral vector
Covishield (Oxford/ AstraZeneca formulation)	Serum Institute of India	49	Feb 15, 2021	Non-replicating viral vector
Spikevax	Moderna	86	Apr 30, 2021	mRNA encoding spike protein
Covilo	Sinopharm (Beijing)	91	May 07, 2021	Inactivated
$Ad26$.COV 2 .S	Janssen	111	Mar 12, 2021	Non-replicating viral vector
CoronaVac	Sinovac	56	Jun 01, 2021	Inactivated
Covaxin	Bharat Biotech	14	Nov 03, 2021	Inactivated
COVOVAX (Novavax formulation)	Serum Institute of India	5	Dec 17, 2021	Recombinant spike protein nanoparticle
Nuvaxovid	Novavax	37	Dec 20, 2021	Recombinant spike protein nanoparticle with adjuvant
Convidecia	CanSino	10	May 19, 2022	Non-replicating viral vector

Table 2.1 Current WHO-approved vaccines

CoronaVac (Sinovac) [\[18](#page-17-9)] within a 6-month time span. After this, four more vaccines were developed which were approved within the next year (Covaxin; Bharat Biotech [[19\]](#page-17-10), COVOVAX; Serum Institute of India [\[20](#page-17-11)], Nuvaxovid; Novavax [[21\]](#page-17-12), and Convidecia; CanSino [[22\]](#page-17-13)). The rapid production of the above vaccines was unprecedented considering that it normally takes at least 10 years from discovery research of a new product through the preclinical, clinical, regulatory approval, manufacturing, and delivery stages [[23–](#page-17-14)[26\]](#page-17-15). However, this was driven by the deadly and disruptive nature of the pandemic and made possible by the unprecedented worldwide cooperation building on existing technologies and with new streamlined approaches to research, development, approval, global manufacturing, and distribution, without sacrifcing testing and safety steps [[27–](#page-17-16)[32\]](#page-18-0).

3 Treatments for COVID-19

The approved drugs for COVID-19 target different aspects of the SARS-CoV-2 infection cycle, for improving COVID-19 disease outcomes. These drugs include (1) monoclonal antibodies that interfere with binding of the receptor binding domain (RBD) of the SARS-CoV-2 spike protein to the ACE2 receptor (a critical step in viral entry into host cells) (Fig. [2.3a\)](#page-5-0); (2) molecular compounds that minimize the

Fig. 2.2 Types of vaccines used as protection against COVID-19 disease. (**a**) mRNA-based vaccine (Comirnaty, SpikeVax). This type of vaccine consists of lipid nanoparticle-encapsulated mRNA molecules encoding a modifed version of the SARS-CoV-2 spike protein. Once injected, this is translated by host immune cells to produce the modifed spike protein molecules which stimulate an adaptive immune response. (**b**) Non-replicating viral vector (Vaxzevria, Covishield, Ad26.COV2.S, Convidecia). This vaccine type consists of a replication-defcient virus carrier containing the full-length DNA coding sequence of the SARS-CoV-2 spike protein which is transcribed into mRNA and then translated into proteins by the host cell to produce an immune response. (**c**) Inactivated vaccine (Covilo, CoronaVac, Covaxin). This type of vaccine contains the whole virus which has been inactivated either by deletion or chemical modifcation of the viral genetic material. (**d**) Recombinant spike protein nanoparticle (also known as a subunit vaccine and a virus-like particle vaccine; COVOVAX). These vaccines resemble virus particles to stimulate an immune response but contain no viral genetic material. (**e**) Recombinant spike protein nanoparticle containing adjuvant (Nuvaxovid). This type of vaccine is a virus-like particle containing an adjuvant to boost the host cell immune response

Fig. 2.3 Types of drugs used for treatment of COVID-19. (**a**) Monoclonal antibodies against spike protein (regdanvimab, casirivimab/imdevimab, sotrovimab, tixagevimab/cilgavimab). These antibody-based treatments disrupt binding of the SARS-CoV-2 RBD to the ACE2 receptor on host cells. (**b**) Anti-infammatory drugs (tocilizumab, anakinra). These drugs block interaction of key cytokines with their receptor signaling cascades and thereby inhibit the hyperactivation of proinfammatory factors involved in the cytokine storm effect. (**c**) Antiviral (remdesivir, PF-07321332/ ritonavir). These drugs inhibit key stages of the viral replication cycle

		Authorization	
Treatment	Institution	granted (date)	Mechanism
Veklury	Gilead Sciences	Jul 03, 2020	Anti-viral: Viral RNA
(Remdesivir)			polymerase inhibitor
Regkirona	Celltrion	Nov 12, 2021	Monoclonal antibody:
(Regdanvimab)			Targeting SARS-CoV-2
			spike protein
Ronapreve	Regeneron	Nov 12, 2021	Monoclonal antibodies:
(Casirivimab/	pharmaceuticals		Targeting SARS-CoV-2
Imdevimab)			spike protein
RoActemra	Hoffmann-La Roche	Dec 07, 2021	Anti-inflammatory:
(tocilizumab)			Monoclonal antibody
			targeting IL-6R
Kineret (Anakinra)	Swedish orphan	Dec 17, 2021	Anti-inflammatory: IL-1R
	Biovitrum		antagonist
Xevudy (sotrovimab)	GlaxoSmithKline and	Dec 17, 2021	Monoclonal antibody:
	Vir biotechnology, Inc.		Targeting SARS-CoV-2
			spike protein
Paxlovid (PF-	Pfizer Inc.	Jan 28, 2022	Anti-viral: 3C-like
07321332/ritonavir)			protease inhibitor
Evusheld	AstraZeneca	Mar 25, 2022	Monoclonal antibodies:
(tixagevimab/			Targeting SARS-CoV-2
cilgavimab)			spike protein

Table 2.2 List of COVID-19 treatments approved for use by the European Medicines Agency

damaging cytokine storm effects of viral infection (Fig. [2.3b](#page-5-0)); and (3) small molecules that prevent proteolytic activation of the SARS-CoV-2 non-structural proteins and replication of the viral RNA (Fig. [2.3c\)](#page-5-0) (Table [2.2\)](#page-6-0). The drugs which have been approved currently for use in either Europe, the United States, or by the World Health Organization (WHO), are indicated below in the order of approval date (earliest to most recent).

3.1 Remdesivir

Remdesivir was the frst antiviral drug to be authorized by the WHO as a treatment for COVID-19. In the United States, the Food and Drug Administration (FDA) approved Remdesivir for emergency use for people greater than 12 years old and heavier than 40 kg (88 lbs) [\[33](#page-18-1)], and it has now been approved for temporary use in more than 50 countries [\[34](#page-18-2)]. It was frst developed in 2016 as an antivirus drug called GS-5734 by Gilead Sciences for the treatment of Ebola virus [\[35](#page-18-3)]. Remdesivir is a nucleotide analogue that inhibits viral RNA synthesis by stalling RNAdependent RNA polymerase complex activity (Fig. [2.3c\)](#page-5-0) [[36\]](#page-18-4). Clinical trials on the use of Remdesivir to improve clinical outcomes in COVID-19 patients have shown mixed results. A meta-analysis conducted by Angamo et al. found that treatment with Remdesivir led to an increase in clinical recovery rate by 21% and 29% on

days 7 and 14, respectively, and the need for supplemental oxygen or mechanical ventilation was reduced by 27% and 47%, respectively, compared to the placebo group [\[37](#page-18-5)]. The same study also found a 39% reduction in mortality on day 14 but with no signifcant difference in this outcome on day 28. One meta-analysis found that 10-day Remdesivir treatment was safe with some adverse effects in hospitalized COVID-19 patients, but there was no reduction in mortality compared to placebo [[38\]](#page-18-6). A more recent meta-analysis of nine randomized controlled trials found no signifcant differences in mortality or use of mechanical ventilation between the Remdesivir and control groups [[39\]](#page-18-7). However, the use of Remdesivir did signifcantly increase recovery ($p = 0.004$) and clinical improvement ($p = 0.020$) rates. Taken together, the results of these studies suggest that further work is required to determine if Remdesivir and related antiviral drugs are effcacious and safe for use in the treatment of COVID-19.

3.2 Anti-Spike Protein Monoclonal Antibodies

One of the most promising therapies in the treatment of COVID-19 disease is the use of monoclonal antibodies that target different epitopes of the spike protein RBD (Fig. [2.3a](#page-5-0)) (Table [2.2\)](#page-6-0).

3.2.1 Regdanvimab

Regdanvimab (originally designated CTP59) was identifed through screening a peripheral blood mononuclear cell library from a convalescent patient as a monoclonal antibody targeting the SARS-CoV-2 spike protein RBD of the viral spike protein [[40\]](#page-18-8). A recent meta-analysis identifed seven studies including 1350 patients in the Regdanvimab arm and 1983 patients in the control group, which showed that Regdanvimab treatment led to decreased mortality and need for supplemental oxygen and/or progression to severe disease outcomes [\[41](#page-18-9)]. However, this did not account for the effects of SARS-CoV-2 variants of concern on the outcomes. It was approved for use in COVID-19 patients with mild or moderate levels of illness by the European Medicines Agency in November 2021.

3.2.2 Casirivimab/Imdevimab Cocktail

Ronapreve (also known as REGN-COV2) is a neutralizing antibody cocktail consisting of Casirivimab and Imdevimab, which target distinct regions of the SARS-Cov-2 spike protein RBD [[42\]](#page-18-10). Theoretically, the antibody cocktail approach may offer advantages over a single monoclonal antibody therapeutic by targeting multiple epitopes and thereby diminishing the chances of immune evasion by emerging SARS-CoV-2 variants. A study of 949 patients with mild-to-moderate COVID-19 who were admitted to hospital during the SARS-CoV-2 Delta wave (July 24 to September 30, 2021) in Fukushima Prefecture, Japan, found that those who received the Casirivimab/Imdevimab cocktail showed signifcantly lower deterioration of symptoms [[43\]](#page-18-11). It was approved for use in COVID-19 patients with mild or moderate levels of illness by the European Medicines Agency on the same date as Regdanvimab (November 12, 2021).

3.2.3 Sotrovimab

Sotrovimab was identifed initially by screening antibodies from a convalescent patient from the SARS-CoV-1 epidemic in 2003 [\[44](#page-18-12)]. This antibody recognizes a conserved epitope in both the SARS-CoV-1 and SARS-CoV-2 spike proteins outside the RBD. This property suggested that this epitope might forestall the mutational escape seen in different SARS-CoV-2 variants [[45\]](#page-18-13). A meta-analysis on the effcacy of different SARS-CoV-2 monoclonal antibody therapies found that Sotrovimab ranked frst by causing a signifcant decrease in the incidence of hospitalization compared to placebo, [\[46](#page-18-14)] and two studies showed that it retained the most activity in neutralizing the Omicron variant [[47,](#page-18-15) [48](#page-18-16)]. Sotrovimab was approved for use by the European Medicines Agency on December 17, 2021 for the treatment of COVID-19 patients over 12 years old and weighing over 40 kg who do not require supplemental oxygen or who have a severe disease risk.

3.2.4 Tixagevimab/Cilgavimab Cocktail

A combination of two monoclonal antibodies, Tixagevimab (also known as AZD8895) and Cilgavimab (AZD1061), was isolated from patients who had recovered from COVID-19 disease [[49\]](#page-18-17). As with the other monoclonal antibody cocktails listed above, Tixagevimab/Cilgavimab binds to non-overlapping epitopes on the spike protein RBD. A trial of 3460 participants who received one dose of this cocktail had a relative risk reduction of 82.8% compared to 1731 individuals who had received placebo [[50\]](#page-18-18). It received approval for medical use for the treatment of COVID-19 in the European Union on March 15, 2022. However, as with the other monoclonal antibody therapeutics, this combination treatment showed a signifcant reduction in effcacy against the Omicron BA.1 and BA.2 SARS-CoV-2 variants [\[46](#page-18-14), [48](#page-18-16)]. This calls to attention the need for new monoclonal antibody therapeutics which target the various Omicron subvariants more effectively.

3.3 Tocilizumab

As the levels of the proinfammatory cytokine interleukin-6 (IL-6) have been found to positively correlate with COVID-19, disease severity and death outcomes drugs which counteract IL-6 signaling might play a role in mitigating these effects [\[51,](#page-18-19) [52\]](#page-19-0).

Tocilizumab is a monoclonal antibody that acts as an IL-6 receptor antagonist and has been approved for the treatment of rheumatoid arthritis, cytokine release syndrome, and other disorders marked by hyper-infammation (Fig. [2.3b](#page-5-0) and Table [2.2](#page-6-0)) [\[53](#page-19-1)]. In a meta-analysis carried out by Maraolo et al., Tocilizumab was associated with higher survival in severe COVID-19 disease patients (odds ratio [OR]: 0.83, 95% confdence interval [CI]: 0.74–0.93), although a larger study size accounting for different dosage regimes will be required to confrm this [\[54](#page-19-2)]. Zhang et al. carried out a meta-analysis of 11 studies consisting of 3406 and 3173 patients assigned to the Tocilizumab and control groups, respectively [\[55](#page-19-3)]. They found that the Tocilizumab group had showed signifcant reductions in the following: 1) the 28–30-day mortality risk, 2) need for mechanical ventilation, 3) time-to-hospital discharge, 4) intensive care unit admission, 5) serious disease trajectory, and 6) serious adverse events, compared to the control group. However, another meta-analysis found that although Tocilizumab signifcantly increased the rate of hospital discharges in COVID-19 patients, it had no effect on all-cause mortality or risk of secondary infections [\[56](#page-19-4)].

Some studies have now been carried out to assess the combined use of Tocilizumab and corticosteroid treatment in COVID-19 patients, and these have generally showed positive effects. Lim et al. carried out a meta-analysis of 13 randomized controlled trials and 24 case-control studies to compare the effcacy of Tocilizumab with corticosteroid treatment on mortality outcomes in 18,702 COVID-19 patients [\[57](#page-19-5)]. This revealed significant reductions in mortality following Tocilizumabdexamethasone (odds ratio [OR]: 0.71, 95% confdence interval [CI]: 0.55–0.92) and Tocilizumab-Methylprednisolone (OR: 0.52, 95% CI: 0.36–0.75) therapies. No reduction in mortality was observed for mono-treatment with Methylprednisolone, and none of the drugs signifcantly reduced the need for mechanical ventilation (OR: 0.72, 95%CI: 0.32–1.60). Hong et al. carried out a retrospective cohort study of 33 COVID-19 patients receiving dexamethasone alone and 33 receiving dexamethasone plus Tocilizumab [[58\]](#page-19-6). This showed that the combination treatment led to a signifcant beneft in a 30-day clinical recovery and reduced the need for supplemental oxygen compared to the dexamethasone mono-treatment group. Furthermore, meta-analysis found that the risk of death for COVID-19 patients treated with a corticosteroid-Tocilizumab combination compared with Tocilizumab alone or placebo control was 26% and 52% lower, respectively [\[59](#page-19-7)]. Considering these promising results, these studies call to attention the need for further testing on the use of COVID-19 treatments targeting different aspects of infammation and immune signaling pathways.

3.4 Anakinra

Considering that hyper-infammation is a key factor in driving severe COVID-19 infections, elevated concentrations of pro-infammatory biomarkers such as interleukin 1 (IL-1) have been identifed in COVID-19 patients who experienced a severe or critically ill outcome (Fig. [2.3b\)](#page-5-0) [[60\]](#page-19-8). Anakinra is a recombinant IL-1 receptor antagonist which has been approved for use in the European Union as an antiinfammatory drug to reduce severity and mortality in COVID-19 patients (Table [2.2](#page-6-0)) [\[61](#page-19-9)]. A meta-analysis which assessed the effects of Anakinra treatment on key infammatory biomarkers found that the serum levels of c-reactive protein (CRP), ferritin, and d-dimer were all reduced in the Anakinra compared to the standard care group [\[62](#page-19-10)]. Another meta-analysis found a signifcant reduction in mortality $(OR = 0.34)$ and need for mechanical ventilation $(OR = 0.68)$ in the Anakinra treatment arm compared with the standard care group [\[63](#page-19-11)]. However, the same study called to attention the need for further studies investigating the safety profle of this drug. These fndings were confrmed by another meta-analysis, although this reported no difference in adverse events between the treatment and standard care groups [[64\]](#page-19-12).

3.5 Ritonavir

Ritonavir was originally developed as an inhibitor of the human immunodefciency virus (HIV) protease [[65,](#page-19-13) [66](#page-19-14)] and has been repurposed for similar use in COVID-19 patients via its ability to inhibit the SARS-CoV-2 3C-like protease enzyme (Fig. [2.3c](#page-5-0) and Table [2.2\)](#page-6-0) [\[67](#page-19-15)]. Thus far, no meta-analyses have demonstrated the effcacy of this compound, either alone or in combination, in preventing serious disease in COVID-19 patients, with several reports of adverse effects [[66,](#page-19-14) [68\]](#page-19-16). We suggest that further studies should be conducted to identify other more efficacious and safer antiviral drug candidates for COVID-19.

4 Effect of SARS-CoV-2 Variants on the Effcacy of Vaccines and Monoclonal Antibody Therapeutics

Although most of the developed vaccines worked well at preventing infections and serious illness courses with the original strain of the virus, most worked less effcaciously against the emerging SARS-CoV-2 variants. Planas et al. tested the sensitivity of Omicron compared to the Delta variant of the WHO-approved monoclonal antibody therapeutics using the S-Fuse assay [\[48](#page-18-16)]. All of these antibodies and antibody mixtures neutralized the Delta variant with IC_{50} concentrations ranging from 16 to 369 ng/mL (Table [2.3](#page-11-0)). However, the Tixagevimab/Cilgavimab combination (Evushield; AstarZeneca) and the Sotrovimab monotherapy (Xevudy; GlaxoSmithKline and Vir Biotechnology, Inc.) showed 85- and three-fold decreases in sensitivity, respectively, against Omicron compared to the Delta variant, and the Casirivimab/Imdevimab combination (Ronapreve; Regeneron) and Regdanvimab (Regkirona; Celltrion) had no detectable neutralizing activity towards the Omicron

	Delta variant (IC_{50} ng/mL)	Omicron variant $(IC_{50}$ ng/mL)
Regdanvimab	92	$9000+$
Casirivimab/Imdevimab	98	$9000+$
Sotrovimab	369	1114
Tixagevimab/Cilgavimab	16	1355

Table 2.3 Sensitivity of omicron compared to delta variant to WHO-approved monoclonal antibody therapeutics. Data taken from Planas et al. [**Planas**]

variant. The same study also tested the potency of antibodies elicited by the Comirnaty (Pfzer/BioNTech) and Vaxzevria (AstraZeneca) vaccines to neutralize the Omicron variant relative to the original SARS-CoV-2 strain and the Delta variant [\[48](#page-18-16)]. For both vaccines, sera were sampled 5 months after a two-dose vaccination schedule. This showed that the neutralizing antibody activity in sera was 3.6-fold lower against the Delta variant compared to the original strain of the Comirnaty vaccine, with no neutralization activity detected against the Omicron variant at the highest concentration. Similarly, the levels of antibodies in sera from Vaxzevria-vaccinated individuals were 2.8-fold lower in the neutralizing the Delta variant compared to the original strain, and no activity was observed against the Omicron variant. Similar fndings were reported by Zhang et al. [[69\]](#page-19-17), Cao et al. [\[70](#page-19-18)], and Carreño et al. [\[71](#page-19-19)]. This underscores the capacity of the Omicron variant to escape the existing therapeutic monoclonal antibody treatments and vaccines.

As a means of predicting the capability of SARS-CoV-2 variants to escape antibody neutralization, Hu et al. developed a computational model to estimate the effect of mutations in the spike protein RBD on antibody neutralization titers [[72\]](#page-19-20). Their results were similar to the experimentally determined neutralization titers of the known variants of concern, and they predicted a 17.4-fold decrease in the susceptibility of Omicron to neutralization.

5 Identifcation of Monoclonal Antibodies and Development of New Vaccines to Overcome the Immune Escape Capabilities of SARS-CoV-2 Variants

5.1 Monoclonal Antibodies

Zakir et al. identifed a broadly neutralizing monoclonal antibody (mAb 9G8) which potently neutralizes the SARS-CoV-2 wild-type, Alpha, and Delta variants [[73\]](#page-20-0). However, this has not been tested with the Omicron variant. A similar result was obtained with mAb 2G1 with respect to neutralizing all SARS-CoV-2 strains, but without testing on the Omicron variant as above [[74\]](#page-20-1). In two in vitro and in vivo studies, Wang et al. found that another monoclonal antibody (mAb 35B5) was capable of neutralizing the original SARS-CoV-2 virus and other variants of concern such as Delta [[75\]](#page-20-2) and Omicron [[76\]](#page-20-3). By using cryo-electron microscopy, they showed that this antibody targets a unique epitope outside the RBD, and this likely disrupts the conformational changes that allow SARS-CoV-2 binding to host ACE2 receptors [[75,](#page-20-2) [76](#page-20-3)]. In a study of 30 healthy volunteers administering a mAb 35B5 nasal spray formulation, it was revealed that nasal mucosal samples collected within 24 h showed effective neutralization against pseudoviruses coated with SARS-CoV-2 spike protein variants including both Delta and Omicron [[77\]](#page-20-4). However, full protection required daily inhalation of the spray, suggesting the need for further studies with optimized formulations to extend the duration of the antibody in the nasal mucosa.

Du et al. identifed a monoclonal antibody (mAb 87G7) with potent in vitro neutralizing activity in vitro against all SARS-CoV-2 variants including the Omicron BA.1/BA.2 subvariants [\[78](#page-20-5)]. Using cryo-electron microscopy and site-directed mutagenesis, they showed that mAb 87G7 targets a conserved hydrophobic amino acid cluster in the ACE2 receptor binding site. Another study isolated two antibodies (EV053273 and EV053286) from convalescent patients after they had been infected with the wild-type version SARS-CoV-2 [\[79](#page-20-6)]. One of these antibodies (EV053273) had potent antiviral activity against wild-type SARS-CoV-2 and the Alpha and Delta variants, and the other (EV053286) had lower activity but neutralized all SARS-CoV-2 variants, including the Omicron BA.1 and BA.2 subvariants. They also found that a combination of these two antibodies blocked infection in vivo using a mouse model. In a similar study, Kovavech et al. identifed a cocktail of two distinct monoclonal antibodies (AX290 and AX677) with high affnity to the SARS-CoV-2 spike protein RBD in all SARS-CoV-2 variants, including Omicron, and administration of this cocktail reduced viral burden and infammation in the lungs of an infected mouse model in vivo [\[80](#page-20-7)]. Finally, another study developed monoclonal antibodies against Omicron and other SARS-CoV-2 variants elicited by vaccination with Convidecia [\[81](#page-20-8)]. One of these antibodies (ZWD12) showed potent neutralization against all strains of concern, including the Omicron variant.

5.2 SARS-CoV-2 Vaccines

5.2.1 Updated Vaccines

From the above fndings, it is clear that the production of new vaccines against the current variant of concern is a pressing matter in gaining control over this pandemic. This includes the production of new vaccines specifcally targeting the Omicron subvariants [\[82](#page-20-9)]. With this objective in mind, a recent study showed that the original Spikevax and Omicron-specifc mRNA vaccines produced by Moderna elicited similar neutralizing responses to the Omicron BA.1 and BA.2 subvariants [[83\]](#page-20-10). However, multiple countries and territories are now faced with outbreaks of Omicron BA.4 and BA.5, which may not be recognized by the time the above vaccines are rolled out. It is also possible that a new variant will branch out from a different part of the SARS-CoV-2 family tree. Thus, most scientists agree that constant updates to

the existing vaccines are essential. Other pharmaceutical companies are testing Omicron-specifc vaccines. For example, Pfzer–BioNTech reported that their new Omicron BA.1-based vaccine produced neutralizing antibody responses against this subvariant that were 2–3 times higher than that seen with a booster dose of their original Comirnaty vaccine [\[84](#page-20-11)]. Another study tested adults who had been doubly vaccinated with Comirnaty and had never tested positive for COVID-19 and then received a booster vaccination with either 1) a third dose of Comirnaty; 2) a recombinant spike protein (MVD614) based on the original SARS-CoV-2 strain or 3) a recombinant spike protein (MVB.1.351) based on the Beta variant [\[85](#page-20-12)]. The results showed that boosting with the MVB.1.351 vaccine resulted in a higher neutralizing antibody response against the original virus as well as the Beta, Delta, and Omicron BA.1 strains, compared to boosting with either the Comirnaty or MVD614 vaccines.

5.2.2 Multivalent Vaccines

One approach that can be taken with vaccines is that of multivalent administrations that simultaneously neutralize multiple variants. This is not a new concept as it has been used for decades with infuenza vaccines each year, such as the simultaneous targeting of different varieties of infuenza A and B strains [[86](#page-20-13)]. It follows that a similar approach could be used to spike RBD sequences from multiple SARS-CoV-2 variants of concern. In line with this objective, Moderna has now developed a bivalent vaccine called mRNA-1273.214, which targets the spike protein of the original SARS-CoV-2 virus as well as the highly mutated Omicron variant [[87\]](#page-20-14). Initial reports from a small trial of 439 participants suggested that this vaccine met the clinical endpoints. The data showed that the mean titer was 2372 for the bivalent vaccine, compared to 1473 for the original Moderna mRNA-1273 vaccine [[88–](#page-20-15)[90\]](#page-20-16). The bivalent vaccine was also well tolerated with a similar side effect profle as the current vaccine. Moderna plans to submit the results of this analysis over the coming weeks to regulators.

6 Natural Products for Improved Management of COVID-19 Patients

Herb-derived natural products have long been used in the management of numerous human ailments since ancient times. With the aid of technical advances in instrumental and biological felds, numerous phytochemicals have been isolated and identifed as active ingredients responsible for the pharmacological actions exerted by famous medicinal plants. With respect to COVID-19, several medicinal plants and phytochemicals have been suggested and explored as potential candidates for the treatment of the disease or alleviation of the symptoms [\[91](#page-21-0)[–93](#page-21-1)]. In fact, herbal medicines have been among the frst options to enter clinical phase testing for COVID-19, owing to their availability and generally good safety and tolerability since most of the medicinal plants have a strong ethnobotanical background of use. From the mechanistic standpoint, phytochemicals might exert protective effects against COVID-19 through several mechanisms, including a direct impact on SARS-CoV-2 replication, and infectivity, regulation of ACE2 receptors and the renin-angiotensin system, anti-infammatory action, and immunomodulatory properties [\[93](#page-21-1), [94](#page-21-2)].

Among the phytochemicals, polyphenols have been the subject of a particular focus for their therapeutic potential in COVID-19 [\[91](#page-21-0)]. As a leading polyphenol, curcumin, the active ingredient of turmeric, has been the subject of several trials in patients at different stages of COVID-19 [\[95](#page-21-3), [96\]](#page-21-4). A systematic review of clinical trials suggested the benefcial effects of different curcuminoid preparations, including nanoformulations and curcumin-piperine combinations, on symptom relief, hospitalization length, and mortality in patients suffering from COVID-19 [\[96](#page-21-4)]. The main mechanism suggested to explain the protective effects of curcumin in COVID-19 is the mitigation of infammatory responses as well as the cytokine storm that is closely associated with end-stage adverse COVID-19 complications [\[95](#page-21-3), [97](#page-21-5), [98](#page-21-6)].

Another herbal product which has shown positive effects in clinical practice is the combination of glycyrrhizin and boswellic acids. Besides anti-infammatory and immunomodulatory activities, both compounds have been reported to exert antiviral effects against SARS-CoV-2 [\[99](#page-21-7), [100\]](#page-21-8). Glycyrrhizin has been proposed to inhibit the main protease (M^{pro}) of SARS-CoV-2, thereby interfering with viral replication [\[101](#page-21-9)]. Additionally, both glycyrrhizin and boswellic acids can interact with the functional spike protein of SARS-CoV-2 and reduce virus infectivity through mitigation of viral entry into the host cells [\[102](#page-21-10), [103\]](#page-21-11). In a randomized, double-blind, and placebo-controlled trial, 50 hospitalized patients with moderate COVID-19 received either the combination of glycyrrhizin (60 mg twice daily) and boswellic acids (200 mg twice daily) or placebo for 14 days [\[104](#page-21-12)]. The fndings revealed a significantly lower rate of mortality in the supplemented $(n = 0)$ vs. placebo $(n = 5)$ group. Moreover, there were signifcant improvements in terms of time to recovery, clinical status, serum CRP levels, and percentage of lymphocytes in the herbal combination group compared with the placebo group.

Chinese herbal medicine (CHM) is a comprehensive system of medicine with a strong ethnobotanical background dating to over 2000 years ago. Since the onset of the pandemic, CHM has been among the frst therapeutic approaches tested for the management of COVID-19. Thus far, numerous herbs and formulae have been studied in patients with COVID-19, and several systematic review has been published [\[105](#page-21-13)[–107](#page-21-14)]. However, the methodological limitations and risk of bias in several of the included trials precluded the possibility of reaching a defnitive judgment on the effcacy and safety of CHM for the management of COVID-19. Recently, a systematic review and meta-analysis of 22 high-quality randomised controlled trials involving 1789 subjects assessed the value of adding CHM to Western medicine in controlling COVID-19 [\[108](#page-21-15)]. The results suggested the safety as well as the beneft of combining CHM with Western medicine in improving clinical, hematological,

and virological indices of COVID-19, particularly in those with mild-to-moderate symptoms [[108\]](#page-21-15). Nevertheless, evidence from long-term and multicenter trials is still required to better clarify the role of CHM in the management of COVID-19.

7 Conclusions and Future Perspectives

The emergence of new highly infective SARS-CoV-2 variants such as Omicron has wreaked havoc around the world by allowing the persistence of a pandemic that has already resulted in considerable damage at the individual, societal, and fnancial levels. Although unprecedented achievements have been made in attempts to stop the spread of COVID-19 disease, the problem has continued due to the mutability of the virus, which renders it with new properties such as increased infectivity and the ability to evade our immune defenses. This review has described efforts aimed at developing new vaccination strategies to keep pace with new SARS-CoV-2 variants as they appear, including variant-specifc and multivalent vaccine designs. This included the use of vaccines that target the spike protein of specifc SARS-CoV-2 strains and multivalent approaches that are directed simultaneously against the original SARS-CoV-2 isolate as well as the Omicron variant. Another possibility is the targeting of other antigenic domains within the virus that lie outside the spike protein RBD, as this may allow the development of a universal coronavirus vaccine [\[109](#page-21-16)].

In addition to the developments in SARS-CoV-2 vaccination strategies, we described pharmaceutical approaches that are currently in use for the treatment of individuals who become ill or suffer from postviral sequelae. Most of the existing drugs consist of either repurposed medicines, such as antiviral compounds and antiinfammatory agents, or monoclonal antibodies obtained from convalescent or vaccinated patients. In addition, other approaches are currently under development to help overcome the limitations of the current methods. In the case of antibody-based therapeutics, one potential strategy is the use of broad coronavirus-directed nanobodies isolated from dromedary camels, which are natural reservoirs of coronaviruses, as these molecules can recognize cavities in proteins that are inaccessible to larger conventional antibodies. With this in mind, Hong et al. constructed a phage display library from camels containing nanobodies capable of protecting transgenic mice-expressing human ACE2 receptors against challenge with the SARS-CoV-2 Beta and Delta variants [[110\]](#page-22-0). In addition, several studies have been conducted which have attempted to identify small molecules that disrupt binding of the SARS-CoV-2 spike protein RBD to the ACE2 receptor. Mediouni et al. screened a library of 15,000 small molecules and identifed a compound called calpeptin, which blocked the entry of some of the SARS'CoV-2 variants in whole cell infectivity assays [\[111](#page-22-1)]. Another study found that an engineered soluble ACE2 peptide had high binding affnity to the spike protein of the original SARS-CoV-2 isolate as well as to the Alpha, Beta, Gamma, and Delta variants [\[112](#page-22-2)]. The same study found that this peptide reduced disease severity and improved survival in a transgenic human

ACE2 mouse model infected with both the original SARS-CoV-2 strain and the Gamma variant. Due to the timing of the above studies, the effects of the SARS-CoV-2, RBD, and ACE2 inhibitors on the Omicron subvariants were not assessed. However, a recent study by Li et al. showed that an engineered ACE2 decoy protein had potent preventative and therapeutic effcacy against both Delta and Omicron in in vivo assays [[113\]](#page-22-3). Finally, we described how several natural products are undergoing preclinical and clinical testing to determine their effcacy as preventative or therapeutic agents to prevent serious outcomes following SARS-CoV-2 infection. The advantage of these approaches is that the molecules concerned generally have good safety profles and are predicted to work across all SARS-CoV-2 variants since they target the effects on the body and not the virus itself.

In conclusion, this review has described the importance of developing vaccines and treatment strategies that keep pace with the new SARS-CoV-2 variants as these emerge. In the case of vaccines and therapeutic antibodies, this could involve the production of broadly neutralizing or variant-specifc products. For treatment approaches, considerable further work is required to identify the most effcacious approaches without the trade-off of poor safety profles. Most of all, it will be important to lay the foundations for a procedural pipeline to cope with the likely appearance of new coronavirus variants.

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